Zhengxiao Zhang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5734985/publications.pdf

Version: 2024-02-01

25 papers

1,253 citations

16 h-index 25 g-index

27 all docs

27 docs citations

times ranked

27

1573 citing authors

#	Article	IF	CITATIONS
1	Precision Microbiome Modulation with Discrete Dietary Fiber Structures Directs Short-Chain Fatty Acid Production. Cell Host and Microbe, 2020, 27, 389-404.e6.	5.1	298
2	Impact of Fecal Microbiota Transplantation on Obesity and Metabolic Syndrome—A Systematic Review. Nutrients, 2019, 11, 2291.	1.7	132
3	Fecal microbial transplantation and fiber supplementation in patients with severe obesity and metabolic syndrome: a randomized double-blind, placebo-controlled phase 2 trial. Nature Medicine, 2021, 27, 1272-1279.	15.2	119
4	Comparison of DNA-, PMA-, and RNA-based 16S rRNA Illumina sequencing for detection of live bacteria in water. Scientific Reports, 2017, 7, 5752.	1.6	116
5	Extraction and modification technology of arabinoxylans from cereal by-products: A critical review. Food Research International, 2014, 65, 423-436.	2.9	102
6	Gut microbiota modulation with long-chain corn bran arabinoxylan in adults with overweight and obesity is linked to an individualized temporal increase in fecal propionate. Microbiome, 2020, 8, 118.	4.9	81
7	Not All Fibers Are Born Equal; Variable Response to Dietary Fiber Subtypes in IBD. Frontiers in Pediatrics, 2020, 8, 620189.	0.9	51
8	Feeding practice influences gut microbiome composition in very low birth weight preterm infants and the association with oxidative stress: A prospective cohort study. Free Radical Biology and Medicine, 2019, 142, 146-154.	1.3	50
9	Prebiotic dietary fibre intervention improves fecal markers related to inflammation in obese patients: results from the Food4Gut randomized placebo-controlled trial. European Journal of Nutrition, 2021, 60, 3159-3170.	1.8	46
10	A Diversified Dietary Pattern Is Associated With a Balanced Gut Microbial Composition of Faecalibacterium and Escherichia/Shigella in Patients With Crohn's Disease in Remission. Journal of Crohn's and Colitis, 2020, 14, 1547-1557.	0.6	43
11	Impact of xylanases on gut microbiota of growing pigs fed corn- or wheat-based diets. Animal Nutrition, 2018, 4, 339-350.	2.1	41
12	Metabolite profiling reveals the interaction of chitin-glucan with the gut microbiota. Gut Microbes, 2020, 12, 1810530.	4.3	31
13	Elucidating the role of the gut microbiota in the physiological effects of dietary fiber. Microbiome, 2022, 10, 77.	4.9	31
14	The role of precision nutrition in the modulation of microbial composition and function in people with inflammatory bowel disease. The Lancet Gastroenterology and Hepatology, 2021, 6, 754-769.	3.7	27
15	Characterization of Nitric Oxide Modulatory Activities of Alkaline-Extracted and Enzymatic-Modified Arabinoxylans from Corn Bran in Cultured Human Monocytes. Journal of Agricultural and Food Chemistry, 2016, 64, 8128-8137.	2.4	20
16	Sex-Specific Differences in the Gut Microbiome in Response to Dietary Fiber Supplementation in IL-10-Deficient Mice. Nutrients, 2020, 12, 2088.	1.7	20
17	Noninvasive monitoring of fibre fermentation in healthy volunteers by analyzing breath volatile metabolites: lessons from the FiberTAG intervention study. Gut Microbes, 2021, 13, 1-16.	4.3	8
18	The impact of epidermal growth factor supernatant on pig performance and ileal microbiotal. Translational Animal Science, 2018, 2, 184-194.	0.4	7

#	Article	lF	Citations
19	Breath volatile metabolome reveals the impact of dietary fibres on the gut microbiota: Proof of concept in healthy volunteers. EBioMedicine, 2022, 80, 104051.	2.7	7
20	Chitin-glucan supplementation improved postprandial metabolism and altered gut microbiota in subjects at cardiometabolic risk in a randomized trial. Scientific Reports, 2022, 12, .	1.6	6
21	Regulation of inducible nitric oxide synthase by arabinoxylans with molecular characterisation from wheat flour in cultured human monocytes. International Journal of Food Science and Technology, 2018, 53, 1294-1302.	1.3	4
22	Leisure-Time Physical Activity before and during Pregnancy Is Associated with Improved Insulin Resistance in Late Pregnancy. International Journal of Environmental Research and Public Health, 2021, 18, 4413.	1.2	4
23	397 Time Series and Correlation Network Analyses to Identify the Role of Maternal Microbiomes on Development of Piglet Gut Microbiome and Susceptibility to Neonatal Porcine Diarrhea Journal of Animal Science, 2018, 96, 213-213.	0.2	2
24	Breath volatile compounds and conjugated polyunsaturated fatty acids as metabolic biomarkers reflecting the interaction between chitin-glucan and the gut microbiota Proceedings of the Nutrition Society, 2020, 79, .	0.4	0
25	Chitin-Glucan Supplementation Altered Gut Microbiota and Improved Postprandial Metabolism in Subjects at Cardiometabolic Risk. Current Developments in Nutrition, 2022, 6, 331.	0.1	0